



SEQUENCE LISTING

<110> KLINEFELTER, Gary

<120> METHOD FOR EVALUATING AND AFFECTING MALE FERTILITY

<130> KLINEFELTER=1C

<140> 09/752,514

<141> 2001-01-03

<150> US 09/123,492

<151> 1998-07-28

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<151> 1998-01-29

<150> US 08/593,677

<151> 1996-01-29

<150> US 60/082,753

<151> 1998-04-23

<160> 4

<170> PatentIn version 3.1

<210> 1

<211> 189

<212> PRT

<213> Homo sapiens

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Met Glu Thr Val Ile Pro Val Asp Val Met Arg Arg Ala Gly Ile Lys
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Val Thr Val Ala Gly Leu Ala Gly Lys Asp Pro Val Gln Cys Ser Arg
35 40 45

Asp Val Val Ile Cys Pro Asp Ala Ser Leu Glu Asp Ala Lys Lys Glu
50 55 60

Gly Pro Tyr Asp Val Val Val Leu Pro Gly Gly Asn Leu Gly Ala Gln
65 70 75 80

Asn Leu Ser Glu Ser Ala Ala Val Lys Glu Ile Leu Lys Glu Gln Glu
85 90 95

Asn Arg Lys Gly Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr Ala Leu
 100 105 110

Leu Ala His Glu Ile Gly Cys Gly Ser Lys Val Thr Thr His Pro Leu
 115 120 125

Ala Lys Asp Lys Met Met Asn Gly Gly His Tyr Thr Tyr Ser Glu Asn
 130 135 140

Arg Val Glu Lys Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro Gly Thr
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Ser Phe Glu Phe Ala Leu Ala Ile Val Glu Ala Leu Asn Gly Lys Glu
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Val Ala Ala Gln Val Lys Ala Pro Leu Val Leu Lys Asp
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 <212> DNA
 <213> Rattus rattus

<220>
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 <222> (190)..(756)
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 cagcagttta tcggaacagc aacaacagca acaaaacctt caaaatttac aagtctttaa 180
 gaaatagaa atg gca tcc aaa aga gct ctg gtc atc cta gcc aaa gga gca 231
 Met Ala Ser Lys Arg Ala Leu Val Ile Leu Ala Lys Gly Ala
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 gag gag atg gag aca gtg att cct gtg gac atc atg cgg cga gct ggg 279
 Glu Glu Met Glu Thr Val Ile Pro Val Asp Ile Met Arg Arg Ala Gly
 15 20 25 30
 att aaa gtc acc gtt gca ggc ttg gct ggg aag gac ccc gtg cag tgt 327
 Ile Lys Val Thr Val Ala Gly Leu Ala Gly Lys Asp Pro Val Gln Cys
 35 40 45
 agc cgt gat gta gtg att tgt ccg gat acc agt ctg gaa gaa gca aaa 375
 Ser Arg Asp Val Val Ile Cys Pro Asp Thr Ser Leu Glu Glu Ala Lys
 50 55 60

aca cag gga cca tac gat gtg gtt gtt ctt cca gga gga aat ctg ggt 423
Thr Gln Gly Pro Tyr Asp Val Val Val Leu Pro Gly Gly Asn Leu Gly
65 70 75

gca cag aac tta tct gag tcg gct ttg gtg aag gag atc ctc aag gag 471
Ala Gln Asn Leu Ser Glu Ser Ala Leu Val Lys Glu Ile Leu Lys Glu
80 85 90

cag gag aac agg aag ggc ctc ata gct gcc atc tgt gcg ggt cct acg 519
Gln Glu Asn Arg Lys Gly Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr
95 100 105 110

gcc ctg ctg gct cac gaa gta ggc ttt gga tgc aag gtt aca tcg cac 567
Ala Leu Leu Ala His Glu Val Gly Phe Gly Cys Lys Val Thr Ser His
115 120 125

cca ttg gct aag gac aaa atg atg aac ggc agt cac tac agc tac tca 615
Pro Leu Ala Lys Asp Lys Met Met Asn Gly Ser His Tyr Ser Tyr Ser
130 135 140

gag agc cgt gtg gag aag gac ggc ctc atc ctc acc agc cgt ggg cct 663
Glu Ser Arg Val Glu Lys Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro
145 150 155

ggg acc agc ttc gag ttt gcg ctg gcc att gtg gag gca ctc agt ggc 711
Gly Thr Ser Phe Glu Phe Ala Leu Ala Ile Val Glu Ala Leu Ser Gly
160 165 170

aag gac atg gct aac caa gtg aag gcc ccg ctt gtt ctc aaa gac 756
Lys Asp Met Ala Asn Gln Val Lys Ala Pro Leu Val Leu Lys Asp
175 180 185

tagagagccc aagccctgga ccctggaccc ccaggctgag caggcattgg aagcccacta 816

gtgtgtccac agcccagtga acctggcatt ggaagcccac tagtgtgtcc acagcccagt 876

gaacctcagg aactaacgtg tgaagtagcc cgctgctcag gaatctcgcc ctggctctgt 936

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<210> 3

<211> 189

<212> PRT

<213> Rattus rattus

<400> 3

Met Ala Ser Lys Arg Ala Leu Val Ile Leu Ala Lys Gly Ala Glu Glu
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Met Glu Thr Val Ile Pro Val Asp Ile Met Arg Arg Ala Gly Ile Lys
20 25 30

Val Thr Val Ala Gly Leu Ala Gly Lys Asp Pro Val Gln Cys Ser Arg
35 40 45

Asp Val Val Ile Cys Pro Asp Thr Ser Leu Glu Glu Ala Lys Thr Gln
50 55 60

Gly Pro Tyr Asp Val Val Val Leu Pro Gly Gly Asn Leu Gly Ala Gln
65 70 75 80

Asn Leu Ser Glu Ser Ala Leu Val Lys Glu Ile Leu Lys Glu Gln Glu
85 90 95

Asn Arg Lys Gly Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr Ala Leu
100 105 110

Leu Ala His Glu Val Gly Phe Gly Cys Lys Val Thr Ser His Pro Leu
115 120 125

Ala Lys Asp Lys Met Met Asn Gly Ser His Tyr Ser Tyr Ser Glu Ser
130 135 140

Arg Val Glu Lys Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro Gly Thr
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Ser Phe Glu Phe Ala Leu Ala Ile Val Glu Ala Leu Ser Gly Lys Asp
165 170 175

Met Ala Asn Gln Val Lys Ala Pro Leu Val Leu Lys Asp
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<220>
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<222> (52)..(618)
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Ser Lys Arg Ala Leu Val Ile Leu Ala Lys Gly Ala Glu Glu Met Glu
5 10 15

aca gtg att cct gtg gac atc atg cgg cga gct ggg att aaa gtc acc	153
Thr Val Ile Pro Val Asp Ile Met Arg Arg Ala Gly Ile Lys Val Thr	
20 25 30	
gtt gca ggc ttg gct ggg aag gac ccc gtg cag tgt agc cgt gat gta	201
Val Ala Gly Leu Ala Gly Lys Asp Pro Val Gln Cys Ser Arg Asp Val	
35 40 45 50	
gtg att tgt ccg gat acc agt ctg gaa gaa gca aaa aca cag gga cca	249
Val Ile Cys Pro Asp Thr Ser Leu Glu Glu Ala Lys Thr Gln Gly Pro	
55 60 65	
tac gat gtg gtt gtt ctt cca gga gga aat ctg ggt gca cag aac tta	297
Tyr Asp Val Val Val Leu Pro Gly Gly Asn Leu Gly Ala Gln Asn Leu	
70 75 80	
tct gag tcg gct ttg gtg aag gag atc ctc aag gag cag gag aac agg	345
Ser Glu Ser Ala Leu Val Lys Glu Ile Leu Lys Glu Gln Glu Asn Arg	
85 90 95	
aag ggc ctc ata gct gcc atc tgt gcg ggt cct acg gcc ctg ctg gct	393
Lys Gly Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr Ala Leu Leu Ala	
100 105 110	
cac gaa gta ggc ttt gga tgc aag gtt aca tcg cac cca ttg gct aag	441
His Glu Val Gly Phe Gly Cys Lys Val Thr Ser His Pro Leu Ala Lys	
115 120 125 130	
gac aaa atg atg aac ggc agt cac tac agc tac tca gag agc cgt gtg	489
Asp Lys Met Met Asn Gly Ser His Tyr Ser Tyr Ser Glu Ser Arg Val	
135 140 145	
gag aag gac ggc ctc atc ctc acc agc cgt ggg cct ggg acc agc ttc	537
Glu Lys Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro Gly Thr Ser Phe	
150 155 160	
gag ttt gcg ctg gcc att gtg gag gca ctc agt ggc aag gac atg gct	585
Glu Phe Ala Leu Ala Ile Val Glu Ala Leu Ser Gly Lys Asp Met Ala	
165 170 175	
aac caa gtg aag gcc ccg ctt gtt ctc aaa gac tagagagccc aagccctgga	638
Asn Gln Val Lys Ala Pro Leu Val Leu Lys Asp	
180 185	
ccctggaccc ccaggctgag caggcattgg aagcccacta gtgtgtccac agcccagtga	698
acctggcatt ggaagcccac tagtgtgtcc acagcccagt gaacctcagg aactaacgtg	758
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